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<10> Renault, Jean-Christophe
Fickensicher, Helmut
Dumoutier, Laure
Hor, Simon

<120> Isolated Cytokine Receptor LICR-2

<130> LUD 5752 NDH

<140> US10/026,106

<141> 2001-12-21

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gtgtacctga catggctcc cagggttggc aacccccagg atgtgaccta tttgtggcc	180
atcagagctc tcccaccgt agacggtggc gcgaagtgga agagtgtgcg ggaaccaagg	240
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gcgtgcggac ggtttctccc agctccaagt cccctgggt ggagtcgaa tacctggatt	360
accttttga agtggagccg gccccacctg tctggtgct caccagacg gaggagatc	420
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#25199012v1<IPT> -Repl Sequence Listing for LUD 5752.wpd

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Glu Ser Glu Tyr Leu Asp Tyr Leu Phe Glu Val Glu Pro Ala Pro Pro			
	115	120	125
Val Leu Val Leu Thr Gln Thr Glu Glu Ile Leu Ser Ala Asn Ala Thr			
	130	135	140
Tyr Gln Leu Pro Pro Cys Met Pro Pro Leu Asp Leu Lys Tyr Glu Val			
	145	150	155
Ala Phe Trp Lys Glu Gly Ala Gly Asn Lys Thr Leu Phe Pro Val Thr			
	165	170	175
Pro His Val Thr Pro His Gly Gln Pro Val Gln Ile Thr Leu Gln Pro			
	180	185	190
Ala Ala Ser Glu His His Cys Leu Ser Ala Arg Thr Ile Tyr Thr Phe			
	195	200	205
Ser Val Pro Lys Tyr Ser Lys Phe Ser Lys Pro Thr Cys Phe Leu Leu			
	210	215	220
Glu Val Pro Glu Ala Asn Trp Ala Phe Leu Val Leu Pro Ser Leu Leu			
	225	230	235
Ile Leu Leu Leu Val Ile Ala Ala Gly Gly Val Ile Trp Lys Thr Leu			
	245	250	255
Met Gly Asn Pro Trp Phe Gln Arg Ala Lys Met Pro Arg Ala Leu Asp			
	260	265	270
Phe Ser Gly His Thr Thr His Pro Val Ala Thr Phe Gln Pro Ser Arg			
	275	280	285
Pro Glu Ser Val Asn Asp Leu Phe Leu Cys Pro Gln Lys Glu Leu Thr			
	290	295	300
Arg Gly Val Arg Pro Thr Pro Arg Val Arg Pro Ala Thr Gln Gln Thr			
	305	310	315
Arg Trp Lys Lys Asp Leu Ala Glu Asp Glu Glu Glu Glu Asp Thr Glu			
	325	330	335
Asp Gly Val Ser Phe Gln Pro Tyr Ile Glu Pro Pro Ser Phe Leu Gly			
	340	345	350
Gln Glu His Gln Ala Pro Gly His Ser Glu Ala Gly Gly Val Asp Ser			
	355	360	365
Gly Arg Pro Arg Ala Pro Leu Val Pro Ser Glu Gly Ser Ser Ala Trp			
	370	375	380
Asp Ser Ser Asp Arg Ser Trp Ala Ser Thr Val Asp Ser Ser Trp Asp			
	385	395	400

Arg Ala Gly Ser Ser Gly Tyr Leu Ala Glu Lys Gly Pro Gly Gln Gly
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 Pro Gly Gly Asp Gly His Gln Glu Ser Leu Pro Pro Pro Glu Phe Ser
 420 425 430
 Lys Asp Ser Gly Phe Leu Glu Glu Leu Pro Glu Asp Asn Leu Ser Ser
 435 440 445
 Trp Ala Thr Trp Gly Thr Leu Pro Pro Glu Pro Pro Asn Leu Val Pro
 450 455 460
 Gly Gly Pro Pro Val Ser Leu Gln Thr Leu Thr Phe Cys Trp Glu Ser
 465 470 475 480
 Ser Pro Glu Glu Glu Glu Glu Ala Arg Glu Ser Glu Ile Glu Asp Ser
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 Asp Ala Gly Ser Trp Gly Ala Glu Ser Thr Gln Arg Thr Glu Asp Arg
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gtgtacctga catggtccc agggcttggc aaccccagg atgtgaccta tttgtggcc	180
tatcagagct ctcccaccg tagacggtgg cgcgaaagtgg aagagtgtgc gggaaccaag	240
gagctgctat gttctatgat gtgcctgaag aaacaggacc tgtacaaca gtcaaggga	300
cgcgtgcgga cggtttctcc cagctccaag tccccctggg tggagtccga atacctggat	360
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ctgagtcca atgccacgta ccagctgccc ccctgcatgc cccactgga tctgaagtat	480
gaggtggcat tctggaagga gggggccgga aacaagacce tattccagt cactcccat	540
ggccagccag tccagatcac tctccagcca gctgccagcg aacaccactg cctcagtgcc	600
agaacctct acacgttcag tgtccgaaa tacagcaagt tctctaagcc cacctgcttc	660
ttgctggagg tcccaggact ttctggaca cacacacct gtggcaacct ttcagcccag	720
cagaccagag tccgtgaatg acttggtcct ctgtcccaa aaggaaactga ccagaggggt	780
caggccgacg cctcgagtca gggccccagc cacccaacag acaagatgga agaaggacct	840

tcagaggac gaagaggagg aggatgagga ggacacagaa gatggcgtca gcttcagcc	900
ctacattgaa ccaccttctt tcctggggca agagcaccag gctccagggc actcggaggc	960
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ttgggattct tcagacagaa gctgggccag cactgtggac tcctcctggg acagggtgg	1080
gtcctctggc tatttggtg agaaggggcc aggccaaggg ccgggtgggg atgggcacca	1140
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ggaagaggag gcgagggaat cagaaattga ggacagcgat gcgggcagct ggggggctga	1380
gagcaccag aggaccgagg acagggggccg gacattgggg cattacatgg ccaggtgagc	1440
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		20					25					30			
Leu	Ser	Gln	Asn	Phe	Ser	Val	Tyr	Leu	Thr	Trp	Leu	Pro	Gly	Leu	Gly
	35					40					45				
Asn	Pro	Gln	Asp	Val	Thr	Tyr	Phe	Val	Ala	Tyr	Gln	Ser	Ser	Pro	Thr
	50				55						60				
Arg	Arg	Arg	Trp	Arg	Glu	Val	Glu	Glu	Cys	Ala	Gly	Thr	Lys	Glu	Leu
65				70					75					80	
Leu	Cys	Ser	Met	Met	Cys	Leu	Lys	Lys	Gln	Asp	Leu	Tyr	Asn	Lys	Phe
		85						90						95	
Lys	Gly	Arg	Val	Arg	Thr	Val	Ser	Pro	Ser	Ser	Lys	Ser	Pro	Trp	Val
		100					105						110		
Glu	Ser	Glu	Tyr	Leu	Asp	Tyr	Leu	Phe	Glu	Val	Glu	Pro	Ala	Pro	Pro
	115						120						125		
Val	Leu	Val	Leu	Thr	Gln	Thr	Glu	Glu	Ile	Leu	Ser	Ala	Asn	Ala	Thr

#25199012v1<IPT> -Repl Sequence Listing for LUD 5752.wpd



130 135 140
 Tyr Gln Leu Pro Pro Cys Met Pro Pro Leu Asp Leu Lys Tyr Glu Val
 145 150 155 160
 Ala Phe Trp Lys Glu Gly Ala Gly Asn Lys Thr Leu Phe Pro Val Thr
 165 170 175
 Pro His Gly Gln Pro Val Gln Ile Thr Leu Gln Pro Ala Ala Ser Glu
 180 185 190
 His His Cys Leu Ser Ala Arg Thr Ile Tyr Thr Phe Ser Val Pro Lys
 195 200 205
 Tyr Ser Lys Phe Ser Lys Pro Thr Cys Phe Leu Leu Glu Val Pro Gly
 210 215 220
 Leu Phe Trp Thr His Thr Pro Cys Gly Asn Leu Ser Ala Gln Gln Thr
 225 230 235 240
 Arg Val Arg Glu

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